

In the Specification:

Please amend the specification as shown:

Please delete the paragraph on page 7, lines 19-21, and replace it with the following paragraph:

Figure 2. Comparison of the amino acid sequence of Gra2 between *N. caninum* (NC) (residues 1-197 of SEQ ID NO: 3) and *T. gondii* (TG) (SEQ ID NO: 13). The unique C-terminal domain of NcGra2 is not shown.

Please delete the paragraph on page 45, lines 16-25, and replace it with the following paragraph:

A BlastP search of the NR Nucleic Acid database with the polypeptide encoded by the 24B cDNA revealed matches to RNA polymerases from a wide variety of taxa. Further inspection revealed these matches were to proline rich regions only, and the matches were otherwise not meaningful. Analysis of the predicted 24B polypeptide showed proline (11.2%), alanine (9%) and serine (8.7%) were the most abundant amino acids present. Further, analysis of the 24B amino acid sequence revealed a high number of short repetitive sequences. Of note are the AYPY (SEQ ID NO: 60) repeats near the carboxy terminus, and the serine/glutamine associated, imperfect repeats in the first half of the polypeptide. The ORF encoding the polypeptide contained just three repeats greater than 7 bases long, and these were AGAGGCA, ACTGGGA and CAATTTCAA.